



## SEQUENCE LISTING

<10> Endress, Gregory A.  
Rosen, Craig A.

<120> Prostate Specific Secreted Protein

<130> PF457

<140> Unassigned

<141> 1999-03-30

<150> 60/080,311

<151> 1998-04-01

<150> 60/080,898

<151> 1998-04-07

<160> 15

<170> PatentIn Ver. 2.0

<210> 1

<211> 825

<212> DNA

<213> Homo sapiens

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<221> CDS

<222> (146)..(682)

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cccagtcaca ggcgagagcc ctggg atg cac cgg cca gag gcc atg ctg ctg 172  
Met His Arg Pro Glu Ala Met Leu Leu  
1 5  
ctg ctc acg ctt gcc ctc ctg ggg ggc ccc acc tgg gca ggg aag atg 220  
Leu Leu Thr Leu Ala Leu Leu Gly Gly Pro Thr Trp Ala Gly Lys Met  
10 15 20 25  
tat ggc cct gga gga ggc aag tat ttc agc acc act gaa gac tac gac 268  
Tyr Gly Pro Gly Gly Gly Lys Tyr Phe Ser Thr Thr Glu Asp Tyr Asp  
30 35 40  
cat gaa atc aca ggg ctg cgg gtg tct gta ggt ctt ctg ctg gtg aaa 316  
His Glu Ile Thr Gly Leu Arg Val Ser Val Gly Leu Leu Leu Val Lys  
45 50 55  
agt gtc cag gtg aaa ctt gga gac tcc tgg gac gtg aaa ctg gga gcc 364  
Ser Val Gln Val Lys Leu Gly Asp Ser Trp Asp Val Lys Leu Gly Ala  
60 65 70  
tta ggt ggg aat acc cag gaa gtc acc ctg cag cca ggc gaa tac atc 412  
Leu Gly Gly Asn Thr Gln Glu Val Thr Leu Gln Pro Gly Glu Tyr Ile  
75 80 85  
aca aaa gtc ttt gtc gcc ttc caa gct ttc ctc cgg ggt gtg gtc atg 460  
Thr Lys Val Phe Val Ala Phe Gln Ala Phe Leu Arg Gly Val Val Met  
90 95 100 105  
tac acc agc aag gac cgc tat ttc tat ttt ggg aag ctt gat ggc cag 508  
Tyr Thr Ser Lys Asp Arg Tyr Phe Tyr Phe Gly Lys Leu Asp Gly Gln  
110 115 120  
atc tcc tct gcc tac ccc agc caa gag ggg cag gtg ctg gtg ggc atc 556  
Ile Ser Ser Ala Tyr Pro Ser Gln Glu Gly Gln Val Leu Val Gly Ile

125                      130                      135  
 tat ggc cag tat caa ctc ctt ggc atc aag agc att ggc ttt gaa tgg 604  
 Tyr Gly Gln Tyr Gln Leu Leu Gly Ile Lys Ser Ile Gly Phe Glu Trp  
       140                      145                      150  
 aat tat cca cta gag gag ccg acc act gag cca cca gtt aat ctc aca 652  
 Asn Tyr Pro Leu Glu Glu Pro Thr Thr Glu Pro Pro Val Asn Leu Thr  
       155                      160                      165  
 tac tca gca aac tca ccc gtg ggt cgc tag ggtggggtat ggggccatcc 702  
 Tyr Ser Ala Asn Ser Pro Val Gly Arg  
       170                      175  
 gagctgaggc catctgggtg gtggtggctg atggtactgg agtaactgag tcgggacgct 762  
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 aaa 825

<210> 2  
 <211> 178  
 <212> PRT  
 <213> Homo sapiens

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 Gly Gly Pro Thr Trp Ala Gly Lys Met Tyr Gly Pro Gly Gly Gly Lys  
                     20                      25                      30  
 Tyr Phe Ser Thr Thr Glu Asp Tyr Asp His Glu Ile Thr Gly Leu Arg  
                     35                      40                      45  
 Val Ser Val Gly Leu Leu Leu Val Lys Ser Val Gln Val Lys Leu Gly  
                     50                      55                      60  
 Asp Ser Trp Asp Val Lys Leu Gly Ala Leu Gly Gly Asn Thr Gln Glu  
                     65                      70                      75                      80  
 Val Thr Leu Gln Pro Gly Glu Tyr Ile Thr Lys Val Phe Val Ala Phe  
                     85                      90                      95  
 Gln Ala Phe Leu Arg Gly Val Val Met Tyr Thr Ser Lys Asp Arg Tyr  
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 Phe Tyr Phe Gly Lys Leu Asp Gly Gln Ile Ser Ser Ala Tyr Pro Ser  
                     115                      120                      125  
 Gln Glu Gly Gln Val Leu Val Gly Ile Tyr Gly Gln Tyr Gln Leu Leu  
                     130                      135                      140  
 Gly Ile Lys Ser Ile Gly Phe Glu Trp Asn Tyr Pro Leu Glu Glu Pro  
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 Thr Thr Glu Pro Pro Val Asn Leu Thr Tyr Ser Ala Asn Ser Pro Val  
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 Gly Arg

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Val Ala Pro Glu Gly Glu Pro Val Thr Gly Ile Trp Ala Ser Leu Lys			
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Asn Asn Ile Leu Ser Ser Ile Arg Leu Lys Phe Gly Asn Asn Trp Ser			
50	55	60	
Gln Glu Tyr Gly Ser Ser Gly Arg Ala Glu Ile Glu Val Lys Leu Asn			
65	70	75	80
Pro Asp Glu Thr Val Leu Gly Phe Ser Gly Ser Phe Tyr Ile Phe Met			
85	90	95	
His Gln Ile Ile Ile Thr Thr Ser Gln Pro Arg Glu Leu Ile Ile Gly			
100	105	110	
Pro Leu Thr Gly Arg Tyr Val Tyr Thr Ser Tyr Pro Glu Asn Pro Asn			
115	120	125	
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 tctcccgga tcttgaggtc acatgcgtgg tgggtggacgt aagccacgaa gaccctgagg 180  
 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240  
 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300  
 ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca acccccatcg 360  
 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc 420  
 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct 480  
 atccaagcga catgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga 540  
 ccacgcctcc cgtgctggac tccgacggct ccttcttct ctacagcaag ctcaccgtgg 600  
 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc 660  
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<210> 5  
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<220>  
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<210> 6  
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<210> 7  
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 <212> DNA  
 <213> Homo sapiens

<400> 7  
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<210> 8  
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 gccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat 180  
 ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt 240  
 ttttggaggc ctaggctttt gcaaaaagct t 271

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<210> 10  
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<400> 11  
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 cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga 180  
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<400> 14  
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<210> 15  
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<400> 15  
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